

Amendments to the Specification:

Please replace paragraph [0149] with the following amended paragraph:

[0149] Any polypeptide sequence encoded by the CA polynucleotide sequences may be analyzed to determine certain preferred regions of the polypeptide. Regions of high antigenicity are determined from data by DNASTAR analysis by choosing values that represent regions of the polypeptide that are likely to be exposed on the surface of the polypeptide in an environment in which antigen recognition may occur in the process of initiation of an immune response. For example, the amino acid sequence of a polypeptide encoded by a CA polynucleotide sequence may be analyzed using the default parameters of the DNASTAR computer algorithm (DNASTAR, Inc., Madison, Wis.; <http://www.dnastar.com/>) world wide web at dnastar.com.